

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q2M2I8	AAK1_HUMAN	Homo sapiens	AP2-associated protein kinase 1	26.955061	T354;T359;T360;T362;S363;T441;T445;S447;T448;T507;S519;S650	S14;T234;S235;T354;T389;T441;T606;S618;T620;S623;S624;S637;S650;T653;T687;S731;S846;S937;S938	34019948;35254053;30620550;37340703;36731797;22564745;35132862;29351928;28657654	MKKFFDSRREQGSGSLGSGSSGGG GTSGLGSGYIGRVFGIGRQQVTVDE VLAEGGFAIVFLVRTSNGMKCALKR MFVNNEHDLQVCKREIQIMRDLSSG HKNIVGYIDSSINNVSSGDVWEVLIL MDFCRGGQVVNLMNQRLQTGFTE NEVLQIFCDTCEAVARLHQCKTPIIH RDLKVENILLHDRGHVLCDFGSAT NKFQNPQTEGVNAVEDEIKKYTTLS YRAPEMVNLYSGKIITTKADIWALGC LLYKLCYFTLPPFGESQVAICDGNFTI PDNSRYSQDMHCLIRYMLEPDPDK RPDIYQVSYFSFKLLKKECPIPNVQN SPIPAKLPEPVKASEAAAKKTQPKAR LTDPIPTTETSIAPRQRPKAGQTQPN PGILPIQPALTPRKRATVQPPQAAG SSNQPGLLASVPQPKPQAPPSPQLP QTQAKQPQAPPTPQQTTPSTQAQGLP AQAQATPQHQQQLFLKQQQQQQQP PPAQQQPAGTFYQQQQAQTQFFQA VHPATQKPAIAQFPVVSQGGSQQL MQNFYQQQQQQQQQQQQQLATA LHQQQLMTQQAALQKQPTMAAGQQ PQPQAAAAPQAPAQEPAIQAPVRQ QPKVQTTPPPQVQKQVGLTPPSSP KTQRAGHRRILSDVTHSAVFGVPAS KSTQLLQAAAAEASLNKSKSATTTPS GSPRTSQQNVYNPSEGSTWNPFD DNFSKLTAEELLNKDFAKLGEGKHP EKLGGSAESLIPGFQSTQGDAFATTS FSAGTAEKRKGGQTVDSGLPLLSVS DPFIPLQVPDAPEKLEGLKSPDTSLL LPDLLPMTDPFGSTSDAVIEKADVAV ESLIPGLEPPVPQRLPSQTESVTSNR TDSLTTGEDSLLDCSLLSNPTDLLEE FAPTAISAPVHKAEDSNLISGFDVP EGSDKVAEDEFDPIPVLITKNPQGGH SRNSSGSSESSLPNLARSLLLVDQLI DL

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